



Sheet 1 of 2

Form PTO-1449 (Rev. 10-92) [modified]							Docket No.: 13020-10	Application No.: 09/618,178		
INFORMATION DISCLOSURE CITATION IN AN APPLICATION							Applicants: Lincoln <i>et al.</i>			
(use several sheets if necessary) #2							Filing Date: 7/18/2000	Group Art Unit: 1653 1653 TECH CENTER 1600/2000		

U.S. PATENT DOCUMENTS												
EXAMINER INITIAL	DOCUMENT NUMBER						DATE	NAME		CLASS	SUB-CLASS	FILING DATE
ccc	4	3	0	9	1	1	2	1/5/1982	Ashley <i>et al.</i>			NOV 15 2000
	4	8	5	2	0	1	7	7/25/1989	Hunkapiller			
	5	2	2	1	5	1	8	6/22/1993	Mills			TECH CENTER 1600/2000
	5	4	1	5	8	3	9	5/16/1995	Zaun <i>et al.</i>			
	5	4	4	3	7	9	1	8/22/1995	Cathcort <i>et al.</i>			
	5	4	4	9	6	2	1	9/12/1995	Klein			
	5	4	5	3	2	4	7	9/26/1995	Beavis <i>et al.</i>			
ccc	5	5	1	6	6	6	3	5/14/1996	Backman <i>et al.</i>			

FOREIGN PATENT DOCUMENTS												
EXAMINER INITIAL	DOCUMENT NUMBER						DATE	COUNTRY	CLASS	SUB-CLASS	TRANSLATION	
											YES	NO
ccc	8	8	0	3	9	5	7	6/2/1988	WIPO			
	9	0	0	4	6	5	1	5/3/1990	WIPO			
	9	2	1	5	7	1	2	9/17/1992	WIPO			
ccc	9	3	1	1	2	6	2	6/10/1993	WIPO			X

OTHER DOCUMENTS
(including Author, Title, Date, Pertinent Pages, Etc.)

ccc	Fisher Scientific Catalog, p. 697, (1983).
	Theo T. Nikiforov <i>et al.</i> , "Genetic Bit Analysis: A Solid Phase Method for Typing Single Nucleotide Polymorphisms," <i>Nucleic Acids Research</i> , vol. 22, No. 20, pp. 4167-4175 (1994).
	David J. Balding <i>et al.</i> , "Statistical Analysis of DNA Fingerprint Data for Ordered Clone Physical Mapping of Human Chromosomes," <i>Bulletin of Mathematical Biology</i> , vol. 53, No. 6, pp. 853-879 (1991).
	Mark Keating, M.D., "Linkage Analysis and Long QT Syndrome Using Genetics to Study Cardiovascular Disease," <i>Circulation</i> , vol. 85, No. 6 pp. 1973-1986 (1992).
ccc	Tosh, Yasunaka <i>et al.</i> , "Determination of Gene Recombination Value," <i>Patent Abstracts of Japan</i> , vol. 018 215 (C-1191) (1994).

EXAMINER: <i>KMCarben</i>	DATE CONSIDERED: <i>12-5-01</i>
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		Filing Date: 7/18/2000	Group Art Unit: 1655 <i>1655</i>

<i>pkp</i>	David C. Mansfield <i>et al.</i> , "Automation of Genetic Linkage Analysis Using Fluorescent Microsatellite Markers," <i>Genomics</i> , vol. 24, pp. 225-233 (1994).
	Mark D. Shriver <i>et al.</i> , "VNTR Allele Frequency Distributions Under the Stepwise Mutation Model: A Computer Simulation Approach," <i>Genetics</i> , vol. 134, pp. 983-993 (1993).
	M. Nata <i>et al.</i> , "Prenatal paternity testing with DNA analyses," <i>International Journal of Legal Medicine</i> , vol. 106, pp. 160-162 (1993).
	Colin P. Kimpton <i>et al.</i> , "Automated DNA Profiling Employing Multiplex Amplification of Short Tandem Repeat Loci," <i>PCR Methods and Applications</i> , vol. 3, pp. 13-22 (1993).
	W. Edward Highsmith, Jr., <i>et al.</i> , "Frequency of the Δ Phe ⁵⁰⁸ Mutation and Correlation with XV.2c/KM-19 Haplotypes in an American Population of Cystic Fibrosis Patients: Result of a Collaborative Study," <i>Clin. Chem.</i> , vol. 36 no 10, pp. 1741-1746 (1990).
<i>Jcc</i>	Andrew G. Clark, "Inference of Haplotypes from PCR-amplified Samples of Diploid Populations," <i>J. Biol. Evol.</i> , vol. 7 no. 2, pp. 111-122 (1990).
	W. Y. Tan, "On the Distribution of the Number of Mutants at the Hypoxanthine-Quanine Phosphoribosomal Transferase Locus in Chinese Hamster Ovary Cells," <i>Mathematical Biosciences</i> , vol. 67, pp. 175-192 (1983).

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